# 71488.sequence.txt SEQUENCE LISTING

<110>	Aston University
<120>	Methods of Producing DNA and Protein Libraries
<130>	W071488PPC
<140>	PCT/GB 03/002573
<141>	2003-06-13
<150>	GB0213816.2
	2002-06-14
<160>	13
<170>	PatentIn version 3.1
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<222> (25)..(27)
<223> n=site of randomisation
<220>
<221> misc_feature
<222> (34)..(36)
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                                                                     52
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<213> Artificial sequence
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<22-3>--PCR primer-
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gactgaagct ttagt
                                                                     15
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     3
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<220>
<223>
      PCR primer
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gtcgctggtc tactac
                                                                     16
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     18
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# 71488.sequence.txt <213> Artificial Sequence <220> <223> partial complementary sequence to SEQ ID 1

<221> misc\_feature

<222> (16)..(18)

<223> nnn represents MAX codon (optimum codon usage for each amino acid
in E. coli)

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18

<210> 5

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<211> 32

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<220> –

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32

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<211> 42

<212> DNA

<213> Artificial Sequence

<220>

<223> consensus zinc finger gene fragment

<220>

<221> misc\_feature

<222> (6)..(8)

<223> n=site of randomisation

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       (24)..(26)
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aatcannntc gctgnnnaat gttnnngtag tcgcatgctg ca.
                                                                     42
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      7
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<223>
       PCR primer
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atgaccatga ttacg
                                                                     15
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<211> 30
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<213> Artificial Sequence
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<223> complementary sequence to SEQ ID 7 and SEQ ID 1 (partially)
<400>
atgaccatga ttacgctatg ccatgactga
                                                                     30
<210> 9
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<211>
       12
<212> DNA
<213> Artificial Sequence
<220>
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       partial complementary sequence to SEQ ID 1
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       nnn represents MAX codon (optimum codon usage for each amino acid
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agctttagtn nn
                                                                        12
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       15
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       PCR primer
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       10
acttgagact gaagc
                                                                       15
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       11
<211>
       15
<212> DNA
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<220>
<223>
     PCR primer
<400> 11
gcatgctaga ctgcc
                                                                       15
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<211> 21
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<213> Artificial Sequence
<220>
<223>
      complementary sequence to SEQ ID 11 and SEQ ID 13 (partially)
<400>
catcagcgta cgatctgacg c
                                                              21
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39